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Table 2 [Genetic and related effects] of [agent] in humans in vivo

End-point	Test	Tissue	Cell type (if specified)	Description of exposed and controls	Response, significance*	Comments	Reference
DNA damage	• DNA adducts • DNA strand breaks • DNA cross-links	Bone marrow, lung, buccal,	Hepatocyte, endothelial, fibroblast,	Industry or occupation; cohort name (e.g. Agricultural Health Study); source of controls; population size;	+ (P < 0.01) (+)		
	DNA oxidationUnscheduled DNA synthesis	lymphocytes , liver, ovary	monocyte, alveolar, mucosa	location	+/- - (-)		
Mutation	OncogeneTumour suppressorOther						
Chromosomal damage	 Chromosomal aberrations Micronucleus formation Sister chromatid exchange Aneuploidy 						
DNA repair							

^{+,} positive

^{–,} negatīve

^{+/-,} equivocal (variable response in several experiments within an adequate study)

^{*} Significance is indicated using asterisks

⁽⁺⁾ or (-), positive/negative in a study of limited quality (specify reason in comments, e.g. only a singe dose tested; data or methods not fully reported; confounding exposures, etc.)

Species, strain, sex	End-point	Test	Tissue	Results	Agent, dose (LED or HID)	Route, duration, dosing regimen	Comments	Reference
Rat F344 male, Mouse B6C3F ₁ female, etc	DNA damage	 DNA adducts DNA strand breaks DNA cross-links DNA oxidation Unscheduled DNA synthesis 	Liver, lung, bone marrow, leukocyte, spermatocytes, oocytes	+ (+) +/- - (-) NT	10	Gavage, inhalation, intraperitoneal injection, drinkingwater; 14 day; ×1, 1×/day, 1×/wk	Address which other doses had a positive response, if dose–response was evident, etc	
	Mutation	 Mouse spot test Mouse specific locus test Dominant lethal test Transgenic animal tests						
	Chromosomal damage	 Chromosomal aberrations Micronucleus formation Sister chromatid exchange Aneuploidy						
	DNA repair							

HID, highest ineffective dose; LED, lowest effective dose (units as reported); NT, not tested

Table 4 [Genetic and related effects] of [agent] in humans and other mammals in vitro [can also be captured in two separate tables]

Species	Tissue, cell line	End-point	Test	Results	Concentration (LEC or HIC)	Comments	Reference
Human	Liver/HepG2, Prostate/LNCaP, Breast/MCF-7, Lung/A549, Cervical/HeLa, Colon CACO-2, Fibroblasts, Lymphocytes, Hepatocytes, other	DNA damage	 DNA adducts DNA strand breaks DNA cross-links DNA oxidation Unscheduled DNA synthesis 	+ (+) +/ - (-) NT			
		Mutation	OncogeneTumour suppressorOther				
		Chromosomal damage	 Chromosomal aberrations Micronucleus formation Sister chromatid exchange Aneuploidy				
		DNA repair					
		Cell transformation					

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Mouse lymphoma L5178Y,	DNA damage	• DNA adducts	+
		DNA strand breaksDNA cross-links	(+)
primary hepatocyte,		• DNA oxidation	+/_
Chinese hamster lung V79,		• Offscheduled DNA synthesis	_
			(-)
hepatocyte, other			NT
	Mutation	• Tk	
		• Ouabain resistance	
		• Other genes	
	Chromosomal damage	Aneuploidy	
		Sister chromatid exchange	
	DNA repair		
	Cell transformation		
	Mouse BALB/c 3T3, Mouse C3H 10T1/2, Mouse primary hepatocyte, Chinese hamster ovary, Chinese hamster lung V79, Syrian hamster, Rat liver RL4, Rat primary	Mouse BALB/c 3T3, Mouse C3H 10T1/2, Mouse primary hepatocyte, Chinese hamster ovary, Chinese hamster lung V79, Syrian hamster, Rat liver RL4, Rat primary hepatocyte, other Mutation Chromosomal damage	Mouse BALB/c 3T3, Mouse C3H 10T1/2, Mouse primary hepatocyte, Chinese hamster ovary, Chinese hamster lung V79, Syrian hamster, Rat liver RL4, Rat primary hepatocyte, other Mutation - Tk - Hprt - Ouabain resistance - Other genes Chromosomal damage Chromosomal damage Chromosomal damage DNA repair

HIC, highest ineffective concentration; LEC, lowest effective concentration, NT, not tested

Table 5 [Genetic and related effects] of [agent] in non-mammalian species in vitro

Phylogenetic	Test system (species, strain)	End-point	Test	Results	Results		Comments	Reference
class				Without metabolic activation	With metabolic activation	concentration(LEC or HIC)		
Acellular systems	DNA, Calf thymus DNA,	DNA damage	• DNA adducts	+	+	As reported		
	plasmid DNA, Prophage		DNA strand breaksDNA cross-links	(+)	(+)			
			IntercalationSOS repair test	+/_	+/_			
				_	_			
				(-)	(-)			
				NT	NT			
Prokaryote (bacteria)	Salmonella typhimurium; TA97, TA98, TA100, TA102, TA104, TA1535, TA1537, TA1538 Escherichia coli; WP2 uvrA	Mutation	• Reverse mutation • Forward mutation					
		DNA repair						
Lower eukaryote (yeast, mould, fungi)	Saccharomyces cerevisiae, Schizosaccharomyces pombe, Aspergillus ridulans;, Neurospora crassa	Mutation	Reverse mutationForward mutationGene conversion					

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		Chromosomal damage	 Intrachromosomal recombination Intragenic mitotic recombination Chromosomal malsegregation 	
Insects	Drosophila melanogaster	Mutation	Somatic mutation and recombination test (SMART) Sex-linked recessive lethal mutations Heritable translocation test Dominant-lethal test	NA
		Chromosomal damage	• Aneuploidy	
		DNA repair		
Plant systems	Vicia faba	DNA damage	• Unscheduled DNA synthesis	D.T.A.
	,,	Divir damage	• Offscheduled DNA synthesis	NA
	, , , , , , , , , , , , , , , , , , , ,	Mutation	Onscheduled DNA syndlesis	NA
			Chromosomal aberrations Micronucleus formation Sister chromatid exchange Aneuploidy	NA

HIC, highest ineffective concentration; LEC, lowest effective concentration, NA, not applicable; NT, not tested